

1 MFKSLT[KV]NKKV[K]PIGENNENEQSSRRNEEGSHPSNQSQQTAAQENK[CEEK][SLKTKSTPV Cng3BFL.pro
1 M-----AKINTQY[SHPSR]-----THLKVKTSDDRLNRAENGLSRAHSSSEETS CNGA3.pro
1 M-----KLS-M[KN]NIINTQ[QS]FVTM-----PNVIMPDIEKEIRMEMENGACSS[SFSE]EDDDSA CNGA1.pro
61 TSE[EH]TNIQDKLSKKNSS[GD]LTNPDPQNAAEPT[GT]VPEQKEMD[FG]KEGPNSPQ[NK]PPA Cng3BFL.pro
44 SVLO[PG]IAMETRGADSGQSGFTGQ-----GIARLS---RLIFL---LRRWAARHVHHQ CNGA3.pro
50 STSE-----ESENENPHARGSEFSYK-----SLRK[GG]PSQ[RE]QYLP[GA]IALFNVN[N]SSNK CNGA1.pro
121 APVIN[EYA]DAQLHNLV[KRM]RQRTALY[KKKL]VEGDLSS[PEA]SPQAKPTAVPPVKE[SD]DRP Cng3BFL.pro
92 DQ[GE]DSFFDRFRGAELKEVSSQ[ESNAQ]ANVGSOEPA[DRGR]SAWPLAKCNTNTSNNTTEEK CNGA3.pro
99 DQ[EE]EE-----K[KKKK]KEKKS[SDD]KNENKNDPEK[KKKK]KDK[EEKKKEEK][SK]DKK CNGA1.pro
181 TEHYRLLWFKV[KRM]PLTEYLKRIKLPNS[DSY]TDRLYLLWILLVTLA---YNWNCWFIPL Cng3BFL.pro
152 KTK-----KKDAIV-----VDP-SSNLYRWLTAT[AL]PVFYNWY---LLTC CNGA3.pro
149 EEE-----KKEVVV-----IDP-SGNTYYNWIFCITLPVMYNWT---MVIA CNGA1.pro
239 RLVEPYQTA[DN]IHFWLIADIIQDIIYLYDMLEIOPRLQ[EV]RGCDIIVDSNELRKHYRTSP Cng3BFL.pro
169 RACFDELOSEYIMLWLVLDYSADVLVVDV-IVRARTGFLEQGLMVSDTNRLWQHYKTT CNGA3.pro
186 RACFDELOSDYLEFWLILLDYVSDIVYLIIDM-EVTRTGYLEQGLLVKEELKLINKYKSNL CHGA1.pro
299 KFOLDVASIIPEDICVIFFFGN-EMFRANRMLKYTSFFEFNHHSIMDKAYIRVIRTT Cng3BFL.pro
248 QFKLDVLSLPTDLAVLKVGCTNYPEVRFNRLKFSRLFEFFDRTETRTNYPNMFRI[GN]LV CNGA3.pro
245 QFKLDVLSLPTDIL[EFK]LGNYPEIRINRLIRFSRMEFFEO[RTET]RTNYPNIFRISNLV CHGA1.pro
358 GYLLE[EL]H[NAC]VY[WAS]NVEGIGTTRWVYDGEKN-----EYLR[CY]WAVR[TL]ITIG Cng3BFL.pro
308 IYILIIHWNACIYFAISKFIFGTD[SWV]PNIS[PEHGR]ISRKYIYSLYWSTLTITIG CNGA3.pro
305 MYIVLIIHWNACVYFYSISKALGF[GN]D[WV]PDINDPEFGRIARKYVYSLYWSTLTITIG CHGA1.pro

FIG. 1.

410 GLEPQTLFEIVFQLLNFFSGVFEVFSLLJQMRDVITGAATANQNYFRACMDPTIAYMNNY Cng3BFL.pro
368 ETPPPVKDEEYLFVVDVFLVGVLFATIVGNVGSISMNNAASRAEFQAKIDSITKOYMQFR CNGA3.pro
365 ETPPPVRDSEYVEFVVDFLIIGVLFATIVGNIGSMISMNNAARAEFOARIIDAIKQYMHFR CNGA1.pro

470 SIPRLVQKRVRTWYETWDSQRMLEDSDILKILLETTVQLALAIADVNESIIISKVDLFKGC D Cng3BFL.pro
428 KVTKDIEIRVIRWFDYLMANKKTVDKEVLSLPDKLKAEIAINVHLDTLKKVRIFQDCE CNGA3.pro
425 NVSKDMEKRVIRWFDYLMANKKTVDKEVLSLPDKLKAEIAINVHLDRLKKVRIFADCE CNGA1.pro

530 TQMIYDMIRIKSVILPGLFVCKKGEIGKEMYIIKHGEVQVLGGPDGTVKVIITLKAGSV Cng3BFL.pro
488 AGLLVELVLKIRFTVESP GDYICKKGDIGKEMYIINEGKLAVVAD-DGVTQFVVLS DGSY CNGA3.pro
485 AGLLVELVLKIQEPQVYSPGDYICKKGDIGREMYIIKEGKLAVVAD-DGVTQFVVLS DGSY CNGA1.pro

590 FGEISILIAAGG---GNRRRTANVVAHGFANILTLDDKKTIQEIVHYPDSERIIMKRR-VL Cng3BFL.pro
547 FGEISILNIKSKSGNRRRTANIRSIGYSDLFCLSKDDLMALTEYPEAKKALEEKGRQIL CNGA3.pro
544 FGEISILNIKSKAGNRRRTANIKSIGYSDLFCLSKDDLMALTEYPDAKIMLEEKGRQIL CNGA1.pro

646 IKQAKATAEATPPRKDLALLFPKREETPKLEKILGGTGKASLARLLKIKREQAQKKEN Cng3BFL.pro
607 MKDNIIDEIIRAGAD-----PKDLEEKV---EQLGSSLDTLQTRFARLL----- CNGA3.pro
604 MKDGLLDLNTANAGSD-----PKDLEEKV---TRMEGSVDLLQTRFARLL-----CNGA1.pro

706 SEGEEEGKENEEDKQKENEDKQKENEEDKQKREPEEKPLDRPECTASPIAVEE Cng3BFL.pro
649 -----AEYNATQMKQRIQSLESQVKG GGDK-----PIADGE CNGA3.pro
646 -----AEYESMOOKLKQRI TKVEKFLKPLIDT-----EFSSIE CNGA1.pro

766 EFHSVRRTVLPRGTSRQSLIIISMAPSAGGEEVLTIEVKEKAKQ. Cng3BFL.pro
682 VP-----GDATKTED-----KQQL. CNGA3.pro
679 GE-----GAESGPID-----ST. CNGA1.pro

FIG. 1. (CONTINUED)



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CCATCCTAATACGACTCACTATAGGGCTCGAGCGGCCGCCGGCGAGGTCATTTCTCTACCTTAAGGCACAGTCATA
AATACAGAGGTTTTCAGAACCACTCAGAGAAGATGTTTAAATCGCTGACAAAAGTCAACAAGGTGAAGCCTATAG
GAGAGAACAATGAGAATGAACAAAAGTTCTCGTCGGAATGAAGAAGGCTCTCACCCAAAGTAATCAGTCTCAGCAAACC
ACAGCACAGGAAGAAAACAAAGGTGAAGAGAAAATCTCTCAAAACCAAGTCAACTCCAGTCACGTCCTGAAGAGCCACA
CACCAACATACAAGACAAACTCTCCAAGAAAAATTCCTCTGGAGATCTGACCACAAACCTTGACCCCTCAAAAATGCAG
CAGAACCAACTGGAACAGTGCCAGAGCAGAGGAAGAAATGGACCCCGGGAAGAAAGGTCCAAAACAGCCCAAAAACAAA
CCGCCGTCAGCTCCTGTTATAATGAGTATGCCGATGCCAGCTACACAACTGGTGAAAAGAAATGCCGTCAAAGAAC
AGCCCTCTACAAGAAAAGTTGGTAGAGGAGATCTCTCCTCACCCGAAGCCAGCCCAAACTGCAAAAGCCCAACGG
CTGTACCACCAAGTAAAAGCGATGATAAGCCAAACAGAACATTACTACAGGCTGTGTGGTTCAAAGTCAAAAAG
ATGCCTTTAAACAGAGTACTTAAAGCGAATTAAACTTCCAAAACAGCATAGATTTCATACACAGATCGACTCTATCTCCT
GTGGCTCTTGCTCACTCTTGCCCTATAACTGGAAGTGTGGTTTATACCACTGCGCCTCGTCTTCCCATATCAAA
CCGCAGACACATACACTACTGGCTTATTGCGGACATCATATGTGATATCATCTACCTTTATGATATGCTATTATC
CAGCCCAGACTCCAGTTTGTAAAGAGGAGAGACATAATAGTGGATTCAAATGAGCTAAGGAACACTACAGGACTTC
TACAAAATTCAGTTGGATGTCGCATCAATAATACCATTTGATATTTGCTACCTCTCTTTGGGTTTAATCCAATGT

FIG. 2.



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TTAGAGCAAATAGGATGTTAAAGTACACTTCATTTTTTGAATTTAATCATCACCTAGAGTCTATAATGGACAAAAGCA
TATATCTACAGAGTTATTCCGAACAACCTGGATACTTGCTGTTTATTTCTGCACATTAATGCCCTGTGTTTATTACTGGGC
TTCAAACATATGAAGGAATTGGCACTACTAGATGGGTGTATGATGGGGAAGGAAACGAGTATCTGAGATGTTATTATT
GGCAGTTCGAACCTTAATTACCAATTGGTGGCCTTCCAGAACCAAACTTTATTGAAAATTGTTTTTCAACTCTTG
AATTTTTTCTGGAGTTTTTGTTCTCCAGTTTAATTGGTCAGATGAGAGATGTGATTGGAGCAGCTACAGCCAA
TCAGAATACTTCCGGCCTGCATGGATGACACCATTTGCCCTACATGAACAATTACTCCATTCCATAAACTTGTGCAAA
AGCGAGTTCGGACTTGGTATGAATATACATGGGACTCTCAAAGAATGCTAGATGAGTCTGATTGCTTAAGACCCCTA
CCAATAAGGTCAGTTAGCCCTCGCCATTGATGTGAACCTTCAGCATCATCAGCAAAGTCGACTTGTTCAAAGGGTTG
TGATACACAGATGATTATGACATGTTGCTAAGATTGAAATCCGTTCTCTATTTGCCCTGGTGACTTTGTCTGCAAAA
AGGAGAAATTGGCAAGGAAATGTATATCATCAAGCATGGAGAAAGTCCAAAGTTCTTGAGGGCCCTGATGGTACTAAA
GTTCTGGTTACTCTGAAAGCTGGTGGTGTGGAGAAATCAGCCTTCTAGCAGCAGGAGGAGAAACCGTCTGAAC
TGCCAATGTGGTGGCCACGGGTTTGCCAATCTTTTAACTCTAGACAAAAAGACCCCTCCAAGAAATCTAGTGCATT
ATCCAGATTCTGAAAGGATCCCTCATGAAGAAAGCCAGAGTGCTTTTAAAGCAGAAAGGCTAAGACCGCAGAAAGCAACC
CCTCCAAGAAAAGATCTTGCCCTCCTCTTCCCACCGAAAGAGACACCCAAACTGTTTAAAACTCTCCTTAGGAGG

FIG. 2. (CONTINUED)



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CACAGGAAAAGCAAGTCTTGCAAGACTACTCAAATTGAAGCGAGAGCAAGCAGCTCAGAAGAAAATTCTGAAG
GAGGAGGGAAGGAAAAGAAAATGAAGATAAACAAAAAGAAAATGAAGATAAACAAAAAGAAAATGAAGATAAAA
GGAAAAGAAAATGAAGATAAAAGATAAAAGGAAGAGAGCCAGAGAGAGCCACTGGACAGACCTGAATGTACAGCAAG
TCCTATTGCAGTGGAGGAAGAACCCCACTCAGTTAGAAGGACAGTTTACCCAGAGGGACTTCTCGTCAATCACTCA
TTATCAGCATGGCTCCTTCTGCTGAGGGCGGAGAGAGGTTCTTACTATTGAAAGTCAAAGAAAAGGCTAAGCAAATAA
ATGTTGATTATCTTTAGATGTGATATAGCTAGTTCCCAAAGTGATTGTACCTAGGATTGTAACCTTAAATTAACGAG
GGGAAACGACATGCTGGGACCCCTTGAGAAACGAAAGGCAAATCCCTAGCTTAGTTTCTAGGACTTATCTGAGAGTGT
GATTTCATGCAGTGGTAATAAGAAGATTATTAAGCAAAAAAAGAAAAAAGAAAAAAGAAAAA

FIG. 2. (CONTINUED)



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ATGTTTAAATCGCTGACAAAAGTCAACAAGGTGAAGCCTATAGGAGAGAAACAATGAGAAATGAACAAAAGTTCTCGTCGGAA
TGAAGAAGGCTCTCACCCAAAGTAATCAGTCTCAGCAAAACCACAGCACAGGAAGAAAACAAGGTGAAGAGAAATCTCTCA
AAACCAAGTCAACTCCAGTCAAGTCTGAAGAGCCACACACCAACATACAAGACAAACTCTCCAAGAAAAATTCCTCTGGA
GATCTGACCACAAACCCTGACCCCTCAAAATGCAGCAGAACCAACTGGAACAGTGCCAGAGCAGAAAGAAATGGACCCCGG
GAAAGAAGGTCCAAACAGCCCAAAAACAACCGCCTGCAGCTCCTGTTATAATGAGTATGCCGATGCCAGCTACACA
ACCTGGTGAAAAGATGCGTCAAAAGAACAGCCCTCTACAAGAAAAAGTTGGTAGAGGGAGATCTCTCTCACCCGAAGCC
AGCCACAACTGCAAAAGCCACGGCTGTACCCAGTAAAGAAAGCGATGATAAGCCAAACAGAACATTACTACAGGCT
GTTGTGGTTCAAAGTCAAAAAGATGCCTTTAACAGAGTACTTAAAGCGAATTAAACTTCCAACAGCATAGATTACATACA
CAGATCGACTCTATCTCCTGTGGCTCTTGCTCACTCTTGCCCTATAACTGGAACCTGCTGGTTATACCACTGCGCCTC
GTCTTCCCATATCAAAACCGCAGACAACATACACTACTGGCTTATTGCGGACATCATATGTGATATCATCTACCTTTATGA
TATGCTATTTATCCAGCCCAAGTCCAGTTTGTAAAGAGGAGGAGACATAATAGTGGAATTCAAATGAGCTAAGGAAACACT
ACAGGACTTCTACAAAATTTAGTTGGATGTGCGCATCAATAATACCATTTTGATATTTGCTACCTCTCTTTGGGTTTAAT
CCAATGTTAGAGCAATAGGATGTTAAAGTACACTTCATTTTTTGAATTTAATCATCACCTAGAGTCTATAATGGACAA
AGCATATATCTACAGAGTTATTCGAACAACTGGATACTTGCTGTTTATTCTGCACATTAATGCCTGTGTTTATTACTGGG

FIG. 3.



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CTTCAAACATATGAAGGAATTGGCACTACTAGATGGGTGTATGATGGGGAAGGAACGAGTATCTGAGATGTTATTATTGG
GCAGTTCGAACTTTAATTACCATTTGGTGGCCTTCCAGAACCCAACTTTATTGAAATTGTTTTTCAACTCTTGAATTT
TTTTTCTGGAGTTTTTGTGTTCTCCAGTTTAATTGGTCAGATGAGAGATGTGATTGGAGCAGCTACAGCCAAATCAGAACT
ACTCCGGCCCTGCAATGGATGACACCAATTGCCCTACATGAACAAATTACTCCATTCTCTAACTTGTGCAAAAAGCGAGTTCCGG
ACTTGGTATGAATATACATGGGACTCTCAAAGAAATGCTAGATGAGTCTGATTTGCTTAAGACCCCTACCAACTACGGTCCA
GTTAGCCCCTCGCCATTGATGTGAACTTCAGCATCATCAGCAAAGTCGACTTGTTCAAAGGTTGTGATACACAGATGATTT
ATGACATGTTGCTAAGATTGAAATCCGTTCTCTATTTGCCCTGGTGACTTTGTCTGCAAAAAGGAGAAATTGGCAAGGAA
ATGTATATCATCAAGCATGGAGAAAGTCCAAGTTCTTGGAGGCCCTGATGGTACTATAAGTTCTGGTTACTCTGAAAGCTGG
GTCCGGTGTGAGAGAAATCAGCCCTTCTAGCAGCAGGAGGAGGAAACCGTCGAACTGCCAAATGTGGTGGCCACGGGTTTG
CCAATCTTTTAACTCTAGACAAAAAGACCCCTCCAAGAAATTCTAGTGCAATTATCCAGATTCTGAAAGGATCCTCATGAAG
AAAGCCAGAGTGCTTTTAAAGCAGAAGGCTAAGACCCGAGAAACCCCTCCAAGAAAAGATCTTGCCCTCCTCTTCCC
ACCGAAAAGAGACACCCAACTGTTTAAACTCTCCTAGGAGGCACAGGAAAAGCAAGTCTTGCAAGACTACTCAAT
TGAAGCGAGAGCAAGCAGCTCAGAAGAAAAGAAAATTCTGAAGGAGGAGAGGAAAAGAAAATGAAGATAAAACA
AAAGAAAATGAAGATAAAACAAAAAGAAAATGAAGATAAAAGGAAAAGAAAATGAAGATAAAAGGAAAGAGAGCCAGA

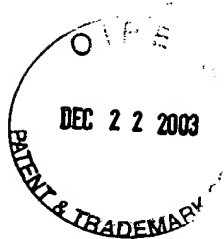
FIG. 3. (CONTINUED)



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AGAGAGCCCACTGGACAGACCTGGAATGTACAGCAAGTCCTATTGCAGTGGAGGAAGAACCCCACTCAGTTAGAAAGGACAG
TTTTACCCAGAGGGACTTCTCGTCAATCACTCATTATCAGCATGGCTCCTTCTGCTGAGGGCGGAGAAAGAGGTTCTTTACT
ATTGAAGTCAAAGAAAAGGCTAAGCAATAA

FIG. 3. (CONTINUED)



MFKSLTKVNKVKPIGENNENEQSSRRNEEGSHPSNQSQQTAAQENKGEKSLKTKSTPVT
EEPHTNIQDKLSKKNSSGDLTTNPDQPQAAEPTGTVPEQKMDPGKEGPNPQNKPAPVI
NEYADAQLHNLVKRMRQRTALYKKLVEGDLSSPEASPTAKPTAVPPVKESDDDKPTEHYR
LLWFKVKKMPLTEYLKRIKLPNSIDSYTDRLYLLWLLVTLAYNWCWFIPRLVFPYQTAD
NIHYWLIADIIICDIIYLYDMLFIOQRLQFVRGGDIIVDSNELRKHRTSTKFFQLDVASIIPF
DICYLFFGFNPMFRANFMLKYTSFFEFNHHLESIMDKAYIYRVIRTTGYLLFILHINACVY
WASNYEGIGTTRWVYDGEENEYLRYYWAVRTLITIGGLPEPQTLFEIVFQLLNFFSGVFVE
SSLIGQMRDVI GAATANQNYFRACMDDTIAYMNNYSIPKL VQKRVRTWYEYTWDSQRM LDES
DLLKTLPTTVQLAL AIDVNF SII SKVDLFKGCDTQMIYDMLLRLLKSVLYLPGDFVCKKGEIG
KEMYIIKHGEVQVLGGPDG TKVLVTLKAGSVFGEISLLAAGGNRRRTANVVAHG FANLLTLD
KKT LQEILVHYPDSE RILMKKARVLLKQAKTAEATPPRKDLALLFP PKEETPKLFKTLILGG
TGKASLARLLKLR EQAAQKKENSEGGEEEGKENEDKOKENEDKOKENEDKDKGR
EPEKPLDRPECTASPIA VEEEPHSVRRTVLP RGT SRQSLIISM APSAEGGEEVLTIEVKEK

AKQ

FIG. 4.